

HNFDL69XX: H.sapiens repeat polymorphism in LIPE gene for hor Sequence Worksheet Human Genome Sciences, Inc.

Sequence Information

Gene Name: H.sapiens repeat polymorphism in LIPE gene to

HGS Code: 563238

Library Catalog: H0271

Sequence ID: HNFDL69XX

Library Name: Human Neutrophil, Activated

Lab Sequenced: HGS

Class: 1

Date Scored:

Date Sequenced:

Lab Scored: HGS

In Group: 18

Group ID: 2514675

Previous Class: 1

Search Results

Overlap	Scor	Overlap Scor Description ,
gb X65642	260	gb x65642 1 260 H.sapiens repeat polymorphism in LIPE gene for hormone sensitive lipase.
gp X54937 HS	93	93 cannabinoid receptor [Homo sapiens] >pir S S17595 cannabinoid receptor - human >sp P21554 CB1R_HUMAN CANNABINOID RECEPTOR 1
gp X55812 RN		Υ.
gp 1.20334 MU		141 EDG-like receptor [Mus musculus]
gp U10699 RN		152 pH218 [Rattus norvegicus] >pir S JC1465 probable G protein-coupled receptor - rat
gp M31210 HUI		210 ECGF1 gene product [Homo sapiens] >pir S A35300 G protein-coupled receptor edg-1 - human >sp P21453 EDG1_HUMAN PROBABLE G P
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Sequence

H H HNFDL69XX unannotated; DNA; 1637 BP.

H. sapiens repeat polymorphism in LIPE gene for hormone sensitive lipase

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Sequence 1637 BP;

CITCATOGGC GCTCTTCACC ecrerceses CCIGGIGAAC CITICCICCIC CGGCCIGGGG CATTGTTCTG GGGGACCCCG GGCACGAGCC TIGCCACCATG CICIGCIGGC TGCTGGCCGC GCTGCTGGGG ATGCTGCCTT TGCTGGGCTG ATTACGATGA GIGCGGCCGG TGGCCGAGAG SOSSISSE GCCCGCACCT CIGGGGGCCA GCCTIGCGG GIGGCCCCCC CACTACAACC CACCCIGCGI CGGGCCTCAG TCAGCCCCCG CCICCACCIT GIGACCIGCT AGICCIGCCA TCCGTCTGGC TCACCAGCCA GECTETCECT **ACTOGGGGG** CACGGGGGG CATGCGGTCG CAGCCIGCIC GCCCGCCCAG GGCCGCCAGC GCIGGCGGG ACAGCIGGG COGGGCCACC GGGGAGGCCA AAGACCAGCC TICACIGCAG TIGGITICCTAC GCCIACCIGG CAACGCIGGG **IGCCIGGIGG** GGITIGCGCIT TGAACGCCAC GCGICTACGG GGAAGGGCT CCAACGIGCI TCTACTATTG TGCTGGAGAA COGAGGATICS ACAGCCGGCT



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GGGATGCATG CCCTGGCAAC GGAAGICCCG GCCCCICICI ааааааааа ааааааа AGAGCACCCT CCATTICCCTA TIGGCTCACCC ACAGGCCCC CCTGGGGTAC CIGAAGIIGC CCGCICCCIC TCACTCCGGA COGGIGICIC CATCATCTAC GIACCIGGG GGGACCACTC CAAGGCCCCC GGCCATCTIC CATCCTCTTC GAACIGCCIG GETGTGGGGG ATGGACTICC AGGAAGCIGI AGICITGOGI AGCITTICGGA GCTTCCACCA COGCTGGGCA TOCTTOOGCA GGCATGGACT TICGGGCIGC IGCIGGCCGA CGICTITIGGC CECCIGCIGA CECCICEICC TECCCCTTTE ACCECTECTC CACCCTTCTE CCCCTCTACT CCAAGCCCTA TIGOCTIGGTICA TOTTTOGCCGG CGTCCTGGCC АГПСААСППС САПСАПОСТА АДАДАДАДАД ДАДДДДДДД GGGCCTCAGT CGAGIGGGIT TIGGACAAGGA CCGGIGGCCI GIGCACGCAA TIGOGGGAGCC CCCACACCIC TIGOGAGGGCC GCAGGGAGGT GIGGAIGGIG GGATCCTGGC AGACGGIGCT AGGCCAGCGG GCAGAAGGCC AGGGCTCCCA GGCTGCAAGG GGTGGACTGT CCCCACAACC COCAACCACC CICGGGGCTT, CIGACGCCAT AIGGACTIGC CCICGCCCIG CAACCACCGG GIGCAGAGCC GATGATCCTG CCTIGTCCAGC TCTGAGGCCA COGGGACTICC CCIGGCCGIC CCGCTTCTGT GTGATTCTGG CCACCICCC TATEGGGAGC GIGCGIGCCA ATCTCCAGCG AGGGACAGCT CIGGCCCGGG GIGCICAGCI TCCTCTGCTG CICAACICGG TOCAACCICI CIGGCCIICI CCACGCCCAG CGGCCCGCC **ACCATCATGG** GTAGGAGCAG AGGGAACGGG GGCAGGCCT TGCGGAGCAT TICECECTC CCGICGAGGC CGGTCAACCC GGGCCCAGGA GCCICIAIGG **IGGIGIGCIG**

Sequence Notes



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HNFDL69XX unannotated; INA; 545 BP.

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Sequence 545 BP;

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EHPGVGASGF LGYRKLCARN LALYGEQGIG HSGASTIDSS AIFRLVQASG GLGALRGLSV YLRGMIWILA FIGLCWLLAA LSCARIFRLA ARAHPASGLS PITPLLCDSG EVPAPLMASV GLEGCKGWIV GCMPWQH*SS IMVKKKKKKK LAVILNSAVNP QKAPRPAARR KARRLLKTVL LRPRDSFRGS AASCLAVLEN LLGMLPLLGW NCLCAFDRCS PAQWFLRKGL QPPGEAMMAT GTPVAPESCQ QAPMOLPGGL SGLLITPYGLA HCLMLITLDKE RSLSFRMREP LFTALAASTF LLVLAAITSH IIYSFRSREV LSSISSVRSI CRAVILSFILCC MILLAFLVCW MRSQRWVYYC SLLPLYSKRY QLAAGGHSRL IVLHYNHSGR LAGRGGPEDC SLLFTAGLRF *SCSLACGWC GPLFGLLLAD CLRLCMRGP AIMVRPVAES LVNITMSDLL ILFCLVIFAG NHRVRARQAL VFGSNILWAQE VLATIMGLYG ATTPPPRRSR GDCLARAVEA GATKTSRVYG TGAAYLANVL

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